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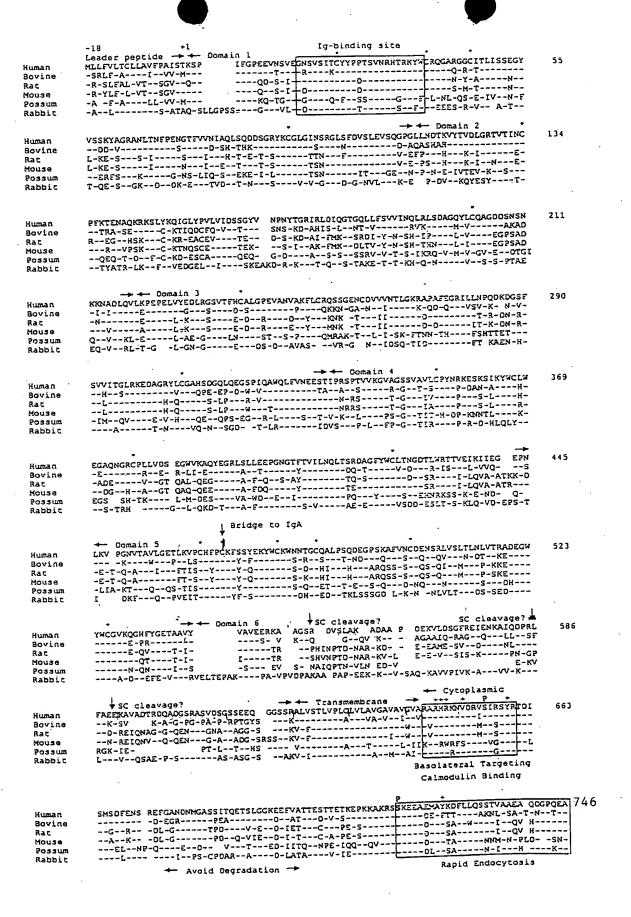
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10	20 	30 -	40 	50 	60
MLLFVLTCLL	AVFPAISTKS	PIFGPEEVNS	VEGNSVSITC	HAMNSIEGTÄ	TRKYWCRQGA
70	80	90	100	110	120
RGGCITLISS	EGYVSSKYAG	RANLTNFPEN	GTFVVNIAQL	SQDDSGRYKC	GLGINSRGLS
130	140	150	160	170	180
FDVSLEVSQG	PGLLNDTXVY	TVDLGRTVTI	NCPFKTENAQ	KSKSLYKQIG	LYPVLVIDSS
190	200	210	220	230	240 I
GYVNBNYTGR 	IRLDIQGTGQ	LLFSVVINQL	RLSDAGQYLC	GYGDD2N2NK	KNADLQVLKP
250	= 260	270	280	290	300
EPELVYEDLR	 GSVTFHCALG	PEVANVAKFL	CRQSSGENCD	VVVNTLGKRA	PAFEGRILLN
310	320	330	340	350	360
PQDXDGSFSV	 VITGLRKEDA	 GRYLCGAHSD	GQLQEGSPIQ	AMOTEANEES	TIPRSPTVVK
370	380	390	400	410	420 I
 GVAGSSVAVL	CPYNRKESKS	IKYWCLWEGA	QNGRCPLLVD	SEGWVKAQYE	GRLSLLEEPG
430	440	450	460	470	480
 NGTFTVILNQ	LTSRDAGFYW	CLTNGDTLWR	 TTVEIKIIEG	EDNTKABGNA I	TAVLGETLKV
490	500	510	520	530	540
PCHFPCKFSS	AEKĀMCKAŅŅ 	TGCQALPSQD	 EGPSKAFVNC	DENSRLVSLT	LNLVTRADEG
sso	· 560	570	530	590	<u> </u>
MAMCGAKÓGH 	 FYGETAAVYV	AVEERKAAGS	RDVSLAKADA	APDEKVLDSG	FREIENKAIQ
610	620	630	640	650	660 I
DPRLFAESKA	VADTRĐQADG	SRASVDSGSS	I EEQGGSSRAL	VSTLVPLGLV	LAVGAVAVGV
670	680	690	700	710	720
 ARARHR <i>K</i> OVD	 RVSIRSYRTD	 ISMSDFENSR	 ADMNDNAGE	SSITQETSLG	GKEEFVATTE
730		750	•	:	
STTETKEPKK	 AKRSSKEEAE	 MAYKDFLLQS	 STVAAEAQDG	PQEA	

Figure 2

7H, 88, 1C

\$ 쑫 æ 보 SF /10H క human pigR stall Ħ 1C, 7H, 6B Rat HUMAN G H FY GETAA V Y V A VEERKAA GSR DVSLA KAOAAP DEK V LOSGFREI ENKAI Q DPRLFAEEKA V A DTRDQADGSRA 8 V D 8 GSSEE Q G G S S R HUMAJGH FYGETAAVYVAVEERKAAGSR DVSLA KADAAPDEKVLDSGFREI ENKAI. RAL GQYYGETTAIYVAVEERTRGSPHNPTDAHA RAKDAPEEEAMESSVREDENKAN. 쿒 RAL GQVYGETTAI YVAVEERTRGSP HNP TOANA RAKOAPEEEAKESSVRED ENKA<u>N WEDDIKUK</u>ADERE I QHAGDQADGNRASGNAGSAGGGSGSSK Humaghfygetaavyvaveerkaagsr ovsla kadaapdekyldsgfrei en<u>kkaluoppkuk</u>apekavadtrdqadgsrasvosgsseeqggssr 쯢 Ŗ B R₂ G Q V Y G ETTAI Y V A V E ERTRGS P MNP T D A NA RAKDA P E E E A NE S S V R E D E N K A N L D P R L F A D E R E I Q N A G D Q A Q E N R A <mark>S G N A G B A G G</mark> Q S G S S K GQVYGETTAI YVAVEERTRGS PHNP TOANA RAKDAPEEEANES S V RED ENKAN<u>LEDDENEREFI</u>ONAGOQAQE H RAS GNAGBAGGOS GS S K GQYYGETTAI YVAVEERTROSPIND TDANA RAKDAPEEEAMESSVRED EKKAN<u>[Edizinius:Xidizi</u>rei ohagdqaqenrasghagsaggosossk GQYYGETTAI YVAVEERTROSPHMP TOANA RAKDAPEEEANESSY, RE<u>IO:EN, YKHYKOD PRIEB</u>ADEREI QNAGOQADEN RASGNAG SAGGQSGSSK GQVYGETTA! YVAVEERTRGSPHMP TDANA RAKDAPEEEAMESSVREDENKAN<u>經的運机过度</u>REI QHAGDQAQENRASGNAGSAGGQSGSSK GQVYGETTAI YVAVEERTRGSPHINY TOANA RAXDAPEEEAMESSYRED ENKANL DP<u>RILEXDEREL</u>QHAGDQAQENRASGNAGGAGGQSGSSK regged c-termini found by Eiffet et al., in SC from human colosbum "B-region" in human colostrum. Intestine B-region is larger end (C-lerminus) of human colosinan SC found by Hughte et al. R=Arg 603 E- G2-938 EKAYADTROQADG9RASVOSGS9EEQGGSSR minimal B-region predicted cleavage site

Figure 3

human pigA staft Redona recognized by a Goal strizerum raised egainst a fusion protein encodita part of domain 5 and all of domain 6 of nat pla? A=ala 568 rayged Clermin found by EDSH et B., human octoburn (EMInt of pl., (1884) Hoppo-Seylor's Z. hysiol, Chem. Bd. 385, S. 1689-1685.) S = Se 570 A=A12516 K=Lys 677 "B-region" in human coloshum. Intestine B-region is larger rrintmai B-region in ivuman colositum end (C-terminus) of human addeterm SC found by Hughes et al. 6-0760 (Hughes at al., (1997), FEBS Letters, A10:443-446.) EXAVAUTRO GADOBRAS V DU GUUE E GOOS BR predicted deavage ena

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Figure 4

CDR 3 FR 4 SFTVNSGYFQHWGQGTLVTVSSGGGGSGGGGSGGGSGGGSEIVLTQSPSTLSASIGDRVTITCRASEGIYHWLA FR 2 CDR 2 FR 2 CDR 2 FR 3 WYQQKPGKAPKLLIYKASSLASGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHYDSTPPT FR 4 FR 5 FR 5 FR 5 FR 6 FR 6 FR 7 FR 6 FR 7 FR 7 FR 8 FR 9 FR 9	GGGSGGGGSEIVL
CDR 3 FR 4 TVNSGYFQHWGQGTLVTVSSGGGGSGGGGGGGGGRR 2 CDR 2 QQKPGKAPKLLIYKASSLASGVPSRFSG	GGSGGGGSEIVLTQSI
CDR.3 FR.4 Links TVNSGYFQHWGQGTLVTVSSGGGGSGGG	nker GGSGGGGSEIVLTQSI
N	
WVROAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCAR	KGRFTISRDNSKNTL
FR 2 CDR 2	
MKYLLPTAAAGLILLAAQPAMADYKAKQVQLVQSGGGLVQPGGSLRLSCAASGFTFSSYAMS	QLVQSGGGLVQPGGSI
PelB leader FLAG	Heavy chain FR 1

determining regions (CDR) of the heavy chain and light chain are indicated. repeated three times), a light chain variable region, a myc epitope tag and a 6HIS tag (for purification by (for secretion in E. coli), a FLAG epitope tag, a heavy chain variable region, a linker sequence (GGGS Immobilized Metal-ion Affinity Chromatography (IMAC)). The framework (FR) and complementarity-The amino acid sequence of the secreted form of the ScFv 4AF is shown. The ScFv consists of a pelb leader